NOV 04 2008

SEQUENCE LISTING

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<110> AFEYAN, NOUBAR B.
      LEE, FRANK D.
      WONG, GORDON G.
      DAS GUPTA, RUCHIRA
      BAYNES, BRIAN
<120> ADZYMES AND USES THEREOF
<130> COTH-P01-001
<140> 10/650,592
<141> 2003-08-27
<150> 60/406,517
<151> 2002-08-27
<150> 60/423,754
<151> 2002-11-05
<150> 60/430,001
<151> 2002-11-27
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Asp Tyr Lys Asp Asp Asp Lys
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Tyr Pro Tyr Asp Val Pro Asp Tyr Ala
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      peptide
Glu Asp Gln Val Asp Pro Arg Leu Ile Asp Gly Lys
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Tyr Thr Asp Ile Glu Met Asn Arg Leu Gly Lys
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Ser Gly Gly Gly
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<400> 9
His Glu Xaa Xaa His
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<400> 10
His His His His His
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1 5

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<211> 26
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<213> Human immunodeficiency virus 1
Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly Arg Lys Lys Arg
                5
                                    10
Arg Gln Arg Arg Pro Pro Gln Gly Ser
            20
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Cys Met His Ile Glu Ser Leu Asp Ser Tyr Thr Cys
                5
                                    10
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      peptide
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Cys Met Tyr Ile Glu Ala Leu Asp Lys Tyr Ala Cys
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<210> 14
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      peptide
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<222> (1)..(1)
<223> A unique residue, such as cysteine or lysine, that
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<221> MOD RES
<222> (2)..(3)
<223> Any residues selected to modulate the affinity of the
      internalizing peptide for different membranes
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<223> see specification as filed for detailed description of
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<400> 14
Xaa Xaa Glu Ala Ala Leu Ala Glu Ala Leu Ala Glu Ala Leu Ala
                                    10
Glu Ala Leu Ala Glu Ala Leu Ala Clu Ala Leu Glu Ala Leu Ala Ala
                                25
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      peptide
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Ala Leu Trp His Trp Trp His
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      peptide
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<223> Thr or Ser
<400> 16
Xaa Trp Leu His Trp Trp Ala
                5
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     peptide
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Gly Gly Gly Ser
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<400> 18
Asp Val Pro Asp Tyr Ala
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Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser
<210> 20
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Gly Gly Val Arg
<210> 21
<211> 639
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Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 5 10 15

Gly Ser Thr Gly Asp Ala Ala Gln Pro Ala Arg Arg Ala Val Arg Ser
20 25 30

Leu Met Thr Ala Thr Ser Glu Tyr Gln Thr Phe Phe Asn Pro Arg Thr 35 40 45

Phe Gly Ser Gly Glu Ala Asp Cys Gly Leu Arg Pro Leu Phe Glu Lys 50 55 60

Lys Ser Leu Glu Asp Lys Thr Glu Arg Glu Leu Leu Glu Ser Tyr Ile
65 70 75 80

Asp Gly Arg Ile Val Glu Gly Ser Asp Ala Glu Ile Gly Met Ser Pro 85 90 95

Trp Gln Val Met Leu Phe Arg Lys Ser Pro Gln Glu Leu Leu Cys Gly
100 105 110

Ala Ser Leu Ile Ser Asp Arg Trp Val Leu Thr Ala Ala His Cys Leu 115 120 125

Leu Tyr Pro Pro Trp Asp Lys Asn Phe Thr Glu Asn Asp Leu Leu Val

Arg Ile Gly Lys His Ser Arg Thr Arg Tyr Glu Arg Asn Ile Glu Lys 145 150 155 160

Ile Ser Met Leu Glu Lys Ile Tyr Ile His Pro Arg Tyr Asn Trp Arg 165 170 175

Glu Asn Leu Asp Arg Asp Ile Ala Leu Met Lys Leu Lys Lys Pro Val 180 185 190

Ala Phe Ser Asp Tyr Ile His Pro Val Cys Leu Pro Asp Arg Glu Thr 195 200 205

Ala Ala Ser 210	Leu Leu	Gln Ala 215	Gly Tyr	Lys Gly	Arg Val	l Thr Gl	y Trp
Gly Asn Leu 225	Lys Glu	Thr Trp 230	Thr Ala	a Asn Val 235		s Gly Gl	n Pro 240
Ser Val Leu	Gln Val 245	Val Asn	Leu Pro	o Ile Val 250	. Glu Arg	g Pro Va 25	_
Lys Asp Ser	Thr Arg 260	Ile Arg	Ile Thi	_	n Met Phe	e Cys Al 270	a Gly
Tyr Lys Pro	-	Gly Lys	Arg Gly 280	/ Asp Ala	Cys Glu 28!	-	p Ser
Gly Gly Pro 290	Phe Val	Met Lys 295	Ser Pro	Phe Asr	a Asn Arg 300	J Trp Ty	r Gln
Met Gly Ile 305	Val Ser	Trp Gly 310	Glu Gly	Cys Asp 315		o Gly Ly	s Tyr 320
Gly Phe Tyr	Thr His	Val Phe	Arg Let	Lys Lys 330	Trp Ile	e Gln Ly 33	
Ile Asp Glr	Phe Gly 340	Glu Gly	Gly Gly 345	-	Gly Gly	Gly Gl 350	y Ser
Gly Gly Gly 355		Met Glu	Val Gli 360	ı Leu Lev	ı Glu Se: 36!	_	y Asp
Leu Val Lys 370	Pro Gly	Gly Ser 375	Leu Lys	s Leu Ser	Cys Ala 380	a Ala Se	r Gly
Phe Thr Phe	Ser Thr	Tyr Gly 390	Met Se	Trp Val	•	n Thr Pr	0 Asp 400
Lys Arg Leu	Glu Trp 405	Val Ala	Thr Ile	e Ser Asr 410	Gly Gly	y Gly Ty 41	
Tyr Tyr Pro	Asp Ser 420	Val Lys	Gly Arg		Ile Se	Arg As	p Asn

Ala Lys Asn Thr Leu Tyr Leu Gln Met Ser Ser Leu Lys Ser Glu Asp 435 440 445

Thr Ala Met Tyr Tyr Cys Ala Arg Arg Glu Arg Tyr Asp Glu Asn Gly 450 455 460

Phe Ala Tyr Trp Gly Arg Gly Thr Leu Val Thr Val Ser Ala Gly Gly 465 470 475 480

Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Gly Ser Asp Ile Val 485 490 495

Met Ser Gln Ser Pro Ser Ser Leu Ala Val Ser Val Gly Glu Lys Ile 500 505 510

Thr Met Ser Cys Lys Ser Ser Gln Ser Leu Phe Asn Ser Gly Lys Gln 515 520 525

Lys Asn Tyr Leu Thr Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys 530 535 540

Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val Pro Asp Arg 545 550 560

Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser 565 570 575

Val Lys Ala Glu Asp Leu Ala Val Tyr Tyr Cys Gln Asn Asp Tyr Ser 580 585 590

His Pro Leu Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg Ala 595 600 605

Asp Ala Ala Pro Thr Ala Arg Gly Gly Pro Glu Gln Lys Leu Ile Ser 610 620

Glu Glu Asp Leu Asn Ser Ala Val Asp His His His His His 625 635

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Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 5 10 15

Gly Ser Thr Gly Asp Ala Ala Gln Pro Ala Arg Arg Ala Val Arg Ser 20 25 30

Leu Met Glu Val Gln Leu Leu Glu Ser Gly Gly Asp Leu Val Lys Pro 35 40 45

Gly Gly Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser 50 55 60

Thr Tyr Gly Met Ser Trp Val Arg Gln Thr Pro Asp Lys Arg Leu Glu 65 70 75 80

Trp Val Ala Thr Ile Ser Asn Gly Gly Gly Tyr Thr Tyr Tyr Pro Asp 85 90 95

Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr 100 105 110

Leu Tyr Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Met Tyr 115 120 125

Tyr Cys Ala Arg Arg Glu Arg Tyr Asp Glu Asn Gly Phe Ala Tyr Trp 130 135 140

Gly Arg Gly Thr Leu Val Thr Val Ser Ala Gly Gly Gly Gly Ser Gly 145 150 155 160

Gly Gly Gly Ser Gly Gly Gly Ser Asp Ile Val Met Ser Gln Ser 165 170 175

Pro Ser Ser Leu Ala Val Ser Val Gly Glu Lys Ile Thr Met Ser Cys 180 185 190

Lys Ser Ser Gln Ser Leu Phe Asn Ser Gly Lys Gln Lys Asn Tyr Leu 195 200 205

Thr	Trp 210	Tyr	Gln	Gln	Lys	Pro 215	Gly	Gln	Ser	Pro	Lys 220	Leu	Leu	Ile	Tyr
Trp 225	Ala	Ser	Thr	Arg	Glu 230	Ser	Gly	Val	Pro	Asp 235	Arg	Phe	Thr	Gly	Ser 240
Gly	Ser	Gly	Thr	Asp 245	Phe	Thr	Leu	Thr	Ile 250	Ser	Ser	Val	Lys	Ala 255	Glu
Asp	Leu	Ala	Val 260	Tyr	Tyr	Cys	Gln	Asn 265	Asp	Tyr	Ser	His	Pro 270	Leu	Thr
Phe	Gly	Gly 275	Gly	Thr	Lys	Leu	Glu 280	Ile	Lys	Arg	Ala	Asp 285	Ala	Ala	Pro
Thr	Gly 290	Gly	Gly	Gly	Ser	Gly 295	Gly	Gly	Gly	Ser	Gly 300	Gly	Gly	Gly	Ser
Met 305	Thr	Ala	Thr	Ser	Glu 310	Tyr	Gln	Thr	Phe	Phe 315	Asn	Pro	Arg	Thr	Phe 320
Gly	Ser	Gly	Glu	Ala 325	Asp	Cys	Gly	Leu	Arg 330	Pro	Leu	Phe	Glu	Lys 335	Lys
Ser	Leu	Glu	Asp 340	Lys	Thr	Glu	Arg	Glu 345	Leu	Leu	Glu	Ser	Tyr 350	Ile	Asp
Gly	Arg	Ile 355	Val	Glu	Gly	Ser	Asp 360	Ala	Glu	Ile	Gly	Met 365	Ser	Pro	Trp
Gln	Val 370	Met	Leu	Phe	Arg	Lys 375	Ser	Pro	Gln	Glu	Leu 380	Leu	Cys	Gly	Ala
Ser 385	Leu	Ile	Ser	Asp	Arg 390	Trp	Val	Leu	Thr	Ala 395	Ala	His	Cys	Leu	Leu 400
Tyr	Pro	Pro	Trp	Asp 405	Lys	Asn	Phe	Thr	Glu 410	Asn	Asp	Leu	Leu	Val 415	Arg
Ile	Gly	Lys	His 420	Ser	Arg	Thr	Arg	Tyr 425	Glu	Arg	Asn	Ile	Glu 430	Lys	Ile
Ser	Met	Leu	Glu	Lys	Ile	Tyr	Ile	His	Pro	Arg	Tyr	Asn	Trp	Arg	Glu

435 440 445

Asn Leu Asp Arg Asp Ile Ala Leu Met Lys Leu Lys Lys Pro Val Ala 450 455 460

Phe Ser Asp Tyr Ile His Pro Val Cys Leu Pro Asp Arg Glu Thr Ala 465 470 475 480

Ala Ser Leu Leu Gln Ala Gly Tyr Lys Gly Arg Val Thr Gly Trp Gly 485 490 495

Asn Leu Lys Glu Thr Trp Thr Ala Asn Val Gly Lys Gly Gln Pro Ser 500 505 510

Val Leu Gln Val Val Asn Leu Pro Ile Val Glu Arg Pro Val Cys Lys 515 520 525

Asp Ser Thr Arg Ile Arg Ile Thr Asp Asn Met Phe Cys Ala Gly Tyr 530 540

Lys Pro Asp Glu Gly Lys Arg Gly Asp Ala Cys Glu Gly Asp Ser Gly 545 550 555 560

Gly Pro Phe Val Met Lys Ser Pro Phe Asn Asn Arg Trp Tyr Gln Met 565 570 575

Gly Ile Val Ser Trp Gly Glu Gly Cys Asp Arg Asp Gly Lys Tyr Gly 580 585 590

Phe Tyr Thr His Val Phe Arg Leu Lys Lys Trp Ile Gln Lys Val Ile 595 600 605

Asp Gln Phe Gly Glu Ala Arg Gly Gly Pro Glu Gln Lys Leu Ile Ser 610 620

Glu Glu Asp Leu Asn Ser Ala Val Asp His His His His His 625 630 635

<210> 23

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<212> DNA

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<223>	Description of Artificial Sequence: Synthetic oligonucleotide	
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ceegga	aaget taatgacege caceagtgag tac	33
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(213)	Artificial Sequence	
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~2237	oligonucleotide	
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ggccc	ctcga gcctctccaa actgatcaat g	31
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	Artificial Sequence	
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tttgga	agagg gaggeggtgg gtetggtggg ggeggtagtg geggaggtgg gageatggag	60

gtgcagctgt tg	72
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cacetecatg eteceacete egecactace gececeacea gacecacege etecetet	cc 60
aaactgatca atg	73
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<213> Artificial Sequence	
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gcaccaactg gaggcggtgg gtctggtggg ggcggtagtg gcggaggtgg gagcatga	cc 60
gccaccagtg agtac	75
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ggtggcggtc atgctcccac ctccgccact accgccccca ccagacccac cgcctcca	gt 60
tggtgcagca tcagc	75
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Tyr Pro Tyr Asp Val Pro Asp Tyr Ala Ser Gly Ser Gly Ser Ser Gly
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Ser Gly Ser Ser Gly Ser Gly Ser Gly Ser Gly Ser Gly Val
Arg
<210> 32
<211> 4
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Ile Thr Pro Arg
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<210> 33
<211> 4
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Ile Thr Leu Arg
<210> 34
<211> 18
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Gly Gly Gly Ser Gly Gly Gly Ser
               5
                                   10
<210> 36
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Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Gly
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Gly Gly Gly Ser
           20
<210> 37
<211> 30
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser
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                               25
<210> 38
<211> 40
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<220>
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Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly
Gly Gly Ser Gly Gly Gly Ser
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<210> 39
<211> 50
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     peptide
<400> 39
Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Gly
               5
Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly
Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly
                           40
Gly Ser
    50
<210> 40
<211> 280
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Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro
               5
                                  10
                                                     15
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- Gly Ser Thr Gly Asp Ile Ala Pro Phe Asp Asp Asp Lys Ile Val 20 25 30
- Gly Gly Tyr Asn Cys Glu Glu Asn Ser Val Pro Tyr Gln Val Ser Leu 35 40 45
- Asn Ser Gly Tyr His Phe Cys Gly Gly Ser Leu Ile Asn Glu Gln Trp 50 55 60
- Val Val Ser Ala Gly His Cys Tyr Lys Ser Arg Ile Gln Val Arg Leu 65 70 75 80
- Gly Glu His Asn Ile Glu Val Leu Glu Gly Asn Glu Gln Phe Ile Asn 85 90 95
- Ala Ala Lys Ile Ile Arg His Pro Gln Tyr Asp Arg Lys Thr Leu Asn 100 105 110
- Asn Asp Ile Met Leu Ile Lys Leu Ser Ser Arg Ala Val Ile Asn Ala 115 120 125
- Arg Val Ser Thr Ile Ser Leu Pro Thr Ala Pro Pro Ala Thr Gly Thr 130 135 140
- Lys Cys Leu Ile Ser Gly Trp Gly Asn Thr Ala Ser Ser Gly Ala Asp 145 150 155 160
- Tyr Pro Asp Glu Leu Gln Cys Leu Asp Ala Pro Val Leu Ser Gln Ala 165 170 175
- Lys Cys Glu Ala Ser Tyr Pro Gly Lys Ile Thr Ser Asn Met Phe Cys 180 185 190
- Val Gly Phe Leu Glu Gly Gly Lys Asp Ser Cys Gln Gly Asp Ser Gly
 195 200 205
- Gly Pro Val Val Cys Asn Gly Gln Leu Gln Gly Val Val Ser Trp Gly 210 215 220
- Asp Gly Cys Ala Gln Lys Asn Lys Pro Gly Val Tyr Thr Lys Val Tyr 225 230 235 240

Asn Tyr Val Lys Trp Ile Lys Asn Thr Ile Ala Ala Asn Ser Thr Arg 245 250 255

Gly Gly Pro Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Ser Ala 260 265 270

Val Asp His His His His His His 275 280

<210> 41

<211> 461

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 41

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro

1 10 15

Gly Ser Thr Gly Asp Ile Ala Pro Phe Asp Asp Asp Lys Ile Val 20 25 30

Gly Gly Tyr Asn Cys Glu Glu Asn Ser Val Pro Tyr Gln Val Ser Leu 35 40 45

Asn Ser Gly Tyr His Phe Cys Gly Gly Ser Leu Ile Asn Glu Gln Trp 50 60

Val Val Ser Ala Gly His Cys Tyr Lys Ser Arg Ile Gln Val Arg Leu 65 70 75 80

Gly Glu His Asn Ile Glu Val Leu Glu Gly Asn Glu Gln Phe Ile Asn 85 90 95

Ala Ala Lys Ile Ile Arg His Pro Gln Tyr Asp Arg Lys Thr Leu Asn 100 105 110

Asn Asp Ile Met Leu Ile Lys Leu Ser Ser Arg Ala Val Ile Asn Ala 115 120 125

Arg Val Ser Thr Ile Ser Leu Pro Thr Ala Pro Pro Ala Thr Gly Thr 130 135 140

Lys Cys Leu 145	Ile Ser	Gly Trp 150	Gly As	sn Thr	Ala Ser 155	Ser Gly	Ala	Asp 160
Tyr Pro Asp	Glu Leu 165	Gln Cys	Leu As	sp Ala 170	Pro Val	Leu Ser	Gln 175	Ala
Lys Cys Glu	Ala Ser 180	Tyr Pro	Gly Ly		Thr Ser	Asn Met 190	Phe	Cys
Val Gly Phe 195	Leu Glu	Gly Gly	Lys As	sp Ser	Cys Gln	Gly Asp 205	Ser	Gly
Gly Pro Val 210	Val Cys	Asn Gly 215	Gln Le	eu Gln	Gly Val 220	Val Ser	Trp	Gly
Asp Gly Cys 225	Ala Gln	Lys Asn 230	Lys Pr	co Gly	Val Tyr 235	Thr Lys	Val	Tyr 240
Asn Tyr Val	Lys Trp 245	Ile Lys	Asn Th	nr Ile 250	Ala Ala	Asn Ser	Leu 255	Val
Pro His Leu	Gly Asp 260	Arg Glu	Lys Ar 26		Ser Val	Cys Pro 270	Gln	Gly
Lys Tyr Ile 275	His Pro	Gln Asn	Asn Se	er Ile	Cys Cys	Thr Lys 285	Cys	His
Lys Gly Thr 290	Tyr Leu	Tyr Asn 295	Asp Cy	/s Pro	Gly Pro 300	Gly Gln	Asp	Thr
Asp Cys Arg 305	Glu Cys	Glu Ser 310	Gly Se	er Phe	Thr Ala 315	Ser Glu	Asn	His 320
Leu Arg His	Cys Leu 325	Ser Cys	Ser Ly	7s Cys 330	Arg Lys	Glu Met	Gly 335	Gln
Val Glu Ile	Ser Ser 340	Cys Thr	Val As	_	Asp Thr	Val Cys 350	Gly	Сув
Arg Lys Asn 355	Gln Tyr	Arg His	Tyr Tr 360	rp Ser	Glu Asn	Leu Phe 365	Gln	Cys

Phe Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His Leu Ser Cys Gln 370 375 380

Glu Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe Leu Arg 385 390 395 400

Glu Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser Leu Glu Cys 405 410 415

Thr Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly Thr Glu Asp 420 425 430

Ser Gly Thr Thr Arg Gly Gly Pro Glu Gln Lys Leu Ile Ser Glu Glu 435 440 445

Asp Leu Asn Ser Ala Val Asp His His His His His 450 455 460

<210> 42

<211> 464

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 42

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 5 10 15

Gly Ser Thr Gly Asp Ile Ala Pro Phe Asp Asp Asp Lys Ile Val

Gly Gly Tyr Asn Cys Glu Glu Asn Ser Val Pro Tyr Gln Val Ser Leu 35 40 45

Asn Ser Gly Tyr His Phe Cys Gly Gly Ser Leu Ile Asn Glu Gln Trp 50 60

Val Val Ser Ala Gly His Cys Tyr Lys Ser Arg Ile Gln Val Arg Leu 65 70 75 80

Gly Glu His Asn Ile Glu Val Leu Glu Gly Asn Glu Gln Phe Ile Asn 85 90 95

- Ala Ala Lys Ile Ile Arg His Pro Gln Tyr Asp Arg Lys Thr Leu Asn 100 105 110
- Asn Asp Ile Met Leu Ile Lys Leu Ser Ser Arg Ala Val Ile Asn Ala 115 120 125
- Arg Val Ser Thr Ile Ser Leu Pro Thr Ala Pro Pro Ala Thr Gly Thr 130 135 140
- Lys Cys Leu Ile Ser Gly Trp Gly Asn Thr Ala Ser Ser Gly Ala Asp 145 150 155 160
- Tyr Pro Asp Glu Leu Gln Cys Leu Asp Ala Pro Val Leu Ser Gln Ala 165 170 175
- Lys Cys Glu Ala Ser Tyr Pro Gly Lys Ile Thr Ser Asn Met Phe Cys 180 185 190
- Val Gly Phe Leu Glu Gly Gly Lys Asp Ser Cys Gln Gly Asp Ser Gly
 195 200 205
- Gly Pro Val Val Cys Asn Gly Gln Leu Gln Gly Val Val Ser Trp Gly 210 215 220
- Asp Gly Cys Ala Gln Lys Asn Lys Pro Gly Val Tyr Thr Lys Val Tyr 225 230 235 240
- Asn Tyr Val Lys Trp Ile Lys Asn Thr Ile Ala Ala Asn Ser Ala Ala 245 250 255
- Ala Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg Asp Ser Val Cys 260 265 270
- Pro Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys Thr 275 280 285
- Lys Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro Gly 290 295 300
- Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala Ser 305 310 315 320

Glu Asn His Leu Arg His Cys Leu Ser Cys Ser Lys Cys Arg Lys Glu 325 330 335

Met Gly Gln Val Glu Ile Ser Ser Cys Thr Val Asp Arg Asp Thr Val 340 345 350

Cys Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr Trp Ser Glu Asn Leu 355 360 365

Phe Gln Cys Phe Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His Leu 370 375 380

Ser Cys Gln Glu Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe 385 390 395 400

Phe Leu Arg Glu Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser 405 410 415

Leu Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly
420 425 430

Thr Glu Asp Ser Gly Thr Thr Arg Gly Gly Pro Glu Gln Lys Leu Ile 435 440 445

Ser Glu Glu Asp Leu Asn Ser Ala Val Asp His His His His His 450 455 460

<210> 43

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<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
 polypeptide

<400> 43

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 5 10 15

Gly Ser Thr Gly Asp Ile Ala Pro Phe Asp Asp Asp Lys Ile Val 20 25 30

Gly Gly Tyr Asn Cys Glu Glu Asn Ser Val Pro Tyr Gln Val Ser Leu 35 40 45

- Asn Ser Gly Tyr His Phe Cys Gly Gly Ser Leu Ile Asn Glu Gln Trp 50 55 60
- Val Val Ser Ala Gly His Cys Tyr Lys Ser Arg Ile Gln Val Arg Leu 65 70 75 80
- Gly Glu His Asn Ile Glu Val Leu Glu Gly Asn Glu Gln Phe Ile Asn 85 90 95
- Ala Ala Lys Ile Ile Arg His Pro Gln Tyr Asp Arg Lys Thr Leu Asn 100 105 110
- Asn Asp Ile Met Leu Ile Lys Leu Ser Ser Arg Ala Val Ile Asn Ala 115 120 125
- Arg Val Ser Thr Ile Ser Leu Pro Thr Ala Pro Pro Ala Thr Gly Thr 130 135 140
- Lys Cys Leu Ile Ser Gly Trp Gly Asn Thr Ala Ser Ser Gly Ala Asp 145 150 155 160
- Tyr Pro Asp Glu Leu Gln Cys Leu Asp Ala Pro Val Leu Ser Gln Ala 165 170 175
- Lys Cys Glu Ala Ser Tyr Pro Gly Lys Ile Thr Ser Asn Met Phe Cys 180 185 190
- Val Gly Phe Leu Glu Gly Gly Lys Asp Ser Cys Gln Gly Asp Ser Gly 195 200 205
- Gly Pro Val Val Cys Asn Gly Gln Leu Gln Gly Val Val Ser Trp Gly 210 215 220
- Asp Gly Cys Ala Gln Lys Asn Lys Pro Gly Val Tyr Thr Lys Val Tyr 225 230 235 240
- Asn Tyr Val Lys Trp Ile Lys Asn Thr Ile Ala Ala Asn Ser Ala Ala 245 250 255
- Ala Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser 260 265 270

Gly Gly Gly Ser Arg Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg Asp Ser Val Cys Pro Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys Thr Lys Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro Gly Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala Ser Glu Asn His Leu Arg His Cys Leu Ser Cys Ser Lys Cys Arg Lys Glu Met Gly Gln Val Glu Ile Ser Ser Cys Thr Val Asp Arg Asp Thr Val Cys Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr Trp Ser Glu Asn Leu Phe Gln Cys Phe Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His Leu Ser Cys Gln Glu Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe Leu Arg Glu Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser Leu Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly Thr Glu Asp Ser Gly Thr Thr Arg Gly Gly Pro

Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Ser Ala Val Asp His

His His His His 485

<211> 239

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 44

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro

1 10 15

Gly Ser Thr Gly Asp Ala Ala Gln Pro Ala Arg Arg Ala Val Arg Ser 20 25 30

Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg Asp Ser Val Cys Pro 35 40 45

Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys Thr Lys
50 55 60

Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro Gly Gln 65 70 75 80

Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala Ser Glu 85 90 95

Asn His Leu Arg His Cys Leu Ser Cys Ser Lys Cys Arg Lys Glu Met 100 105 110

Gly Gln Val Glu Ile Ser Ser Cys Thr Val Asp Arg Asp Thr Val Cys
115 120 125

Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr Trp Ser Glu Asn Leu Phe 130 135 140

Gln Cys Phe Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His Leu Ser 145 150 155 160

Cys Gln Glu Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe 165 170 175

Leu Arg Glu Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser Leu 180 185 190 Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly Thr 195 200 205

Glu Asp Ser Gly Thr Thr Arg Gly Gly Pro Glu Gln Lys Leu Ile Ser 210 $\,$ 215 $\,$ 220 $\,$

Glu Glu Asp Leu Asn Ser Ala Val Asp His His His His His 225 230 235

<210> 45

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic peptide

<400> 45

Arg Pro Leu Ala Leu Trp Arg Ser